
Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Fri Jul 06 13:55:53 EDT 2007

Validated By CRFValidator v 1.0.2

Application No: 10529000 Version No: 1.1

Input Set:

Output Set:

Started: 2007-07-06 13:55:30.902

Finished: 2007-07-06 13:55:32.188

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 286 ms

Total Warnings: 52

Total Errors: 0

No. of SeqIDs Defined: 55

Actual SeqID Count: 55

Error code		Error Descripti	ion								
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(4)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(5)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(6)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(7)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(8)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(9)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(10)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(11)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(12)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(13)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(14)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(15)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(16)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(17)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(18)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(19)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(20)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(21)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(22)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(23)

Input Set:

Output Set:

Started: 2007-07-06 13:55:30.902 **Finished:** 2007-07-06 13:55:32.188

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 286 ms

Total Warnings: 52

Total Errors: 0

No. of SeqIDs Defined: 55

o. of SeqIDs Defined: 55
Actual SeqID Count: 55

Error code Error Description

This error has occured more than 20 times, will not be displayed

55

60

65

70

```
<110> SATO, Noriyuki
     TSUKAHARA, Tomohide
     NABETA, Yuki
     KAWAGUCHI, Satoshi
     IKEDA, Hideyuki
     WADA, Takuro
     YAMASHITA, Toshihiko
<120> TUMOR ANTIGEN PROTEIN AND USE THEREOF
<130> 0020-5360PUS1
<140>
      US 10/529,000
<141>
      2005-03-24
<150> PCT/JP03/12037
<151> 2003-09-19
<150> JP 2002-282345
<151> 2002-09-27
<160> 55
<170> PatentIn Ver. 2.1
<210> 1
<211> 2068
<212> DNA
<213> Homo sapiens
<220>
<221> CDS
<222> (337)..(1878)
<400> 1
ctgcacaggg agtttgtctt gtgcaaacaa tttccaaggc agcgttttct tccctgcctg 60
ggagtgcagg gctcagcgcc ttcactttgg aactgactca gagacctaaa gaagcccacc 120
tggccagcgg gaaggggggc cgccgccgc tcccggtttt gggcagccct ggccagctcc 180
ctgtggcctt ggaggacttc caccgggcag gcgttcccat gatgccaggc taccaggcgc 240
gggggattcc tgcaggccgg cgctgctttt cttagaaccc cctttctaga aaagtacacc 300
tggaggtttt gcttcaaaga gaggagaggc agcagc atg gcg agt gtc ctg tcc
                                        Met Ala Ser Val Leu Ser
cga cgc ctt gga aag cgg tcc ctc ctg gga gcc cgg gtg ttg gga ccc
                                                                   402
Arg Arg Leu Gly Lys Arg Ser Leu Leu Gly Ala Arg Val Leu Gly Pro
             10
                                 15
agt gcc tcg gag ggg ccc tcg gct gcc cca ccc tcg gag cca ctg cta
                                                                   450
Ser Ala Ser Glu Gly Pro Ser Ala Ala Pro Pro Ser Glu Pro Leu Leu
        25
                             30
gaa ggg gcc gct ccc cag cct ttc acc acc tct gat gac acc ccc tgc
Glu Gly Ala Ala Pro Gln Pro Phe Thr Thr Ser Asp Asp Thr Pro Cys
     40
                         45
cag gag cag ccc aag gaa gtc ctt aag gct ccc agc acc tcg ggc ctt
                                                                   546
Gln Glu Gln Pro Lys Glu Val Leu Lys Ala Pro Ser Thr Ser Gly Leu
```

_	_		-		_			_	_	-	tat Tyr					594
GIII	GIII	vai	АІА	75	GIII	FIO	GIY	GIII	80	vai	TYL	vai	пр	85	Gly	
			_			_			_		agc		_			642
Gly	Gln	Glu	Cys 90	Thr	Gly	Leu	Val	Glu 95	Gln	His	Ser	Trp	Met 100	Glu	Gly	
caq	ata	acc		taa	cta	cta	aaa		aaσ	cta	cag	atc		tac	agg	690
_			-		_	_		_	_	_	Gln	-	_	_		
		105					110					115				
gtg	gag	gag	gtg	tgg	ctg	gca	gag	ctg	cag	ggc	ccc	tgt	ccc	cag	gca	738
Val		Glu	Val	Trp	Leu		Glu	Leu	Gln	Gly	Pro	Cys	Pro	Gln	Ala	
cca	120	cta	nan	CCC	ana	125	cad	acc	c+ a	acc	130 tac	nns	aaa	ata	tcc	786
		_	-			_	_	_	_	-	Tyr			-		, 00
135					140					145	_	_			150	
agg	aac	atc	gat	gtc	cca	aag	agg	aag	tcg	gac	gca	gtg	gaa	atg	gat	834
Arg	Asn	Ile	Asp		Pro	Lys	Arg	Lys		Asp	Ala	Val	Glu		Asp	
~~~	2 t &	2+4	aaa	155	2+4	a+ a	at a	2 9 9	160	at a	+ a a	+ ~ ~	200	165	~++	882
	_	_		_	_		_	_		_	tcc Ser	_	_		_	002
			170					175				- 1	180			
gta	cag	agt	cct	ccc	aaa	acc	gag	gcc	aac	ttc	tct	gct	tcc	cgt	gcg	930
Val	Gln		Pro	Pro	Gly	Thr	Glu	Ala	Asn	Phe	Ser	Ala	Ser	Arg	Ala	
		185					190					195				070
										_	tcg Ser					978
AIA	200	Asb	FIO	ттр	пуз	205	Ser	Gly	Asp	116	210	Asp	Set	Gry	Ser	
agc	act	acc	agc	ggt	cac	tgg	agt	ggg	agc	agt	ggt	gtc	tcc	acc	CCC	1026
Ser	Thr	Thr	Ser	Gly	His	Trp	Ser	Gly	Ser	Ser	Gly	Val	Ser	Thr	Pro	
215					220					225					230	
					_	-	_		_		ttg		-	-		1074
ser	Pro	Pro	HIS	235	GIN	Ala	ser	Pro	Lуs 240	ıyr	Leu	GIĀ	Asp	245	Pne	
ggt	tct	CCC	caa		gat	cat	ggc	ttt		acc	gat	cct	gac		ttc	1122
Gly	Ser	Pro	Gln	Thr	Asp	His	Gly	Phe	Glu	Thr	Asp	Pro	Asp	Pro	Phe	
			250					255					260			
_	_	-	-		-		-		-	_	aac			_		1170
Leu	Leu	265	GIU	Pro	Ala	Pro	Arg 270	гуз	Arg	гуз	Asn	275	Val	гуѕ	Val	
atg	tac		tgc	ctg	tgg	cca		tgt	ggc	aaa	gtt		cgc	tcc	att	1218
Met	Tyr	Lys	Cys	Leu	Trp	Pro	Asn	Суз	Gly	Lys	Val	Leu	Arg	Ser	Ile	
	280					285					290					
				-		-		-			ctg		-			1266
Val 295	GIY	Ile	Lys	Arg	His	Val	Lys	Ala	Leu	His	Leu	GLY	Asp	Thr	Val 310	
	tct	gat	caq	ttc		caa	gag	gag	gat		tac	tac	aca	gag		1314
_		_	_		_				-		Tyr					
				315					320					325		
cag	ctg	aag	gag	gaa	tct	gct	gct	gct	gct	gct	gct	gct	gcc	gca	ggc	1362
Gln	Leu	Lys		Glu	Ser	Ala	Ala		Ala	Ala	Ala	Ala		Ala	Gly	
acc	cca	at c	330 cct	aaa	act	CCC	acc	335 t.cc	gan	cca	gct	aac	340 acc	aaa	age	1410
		-									Ala				-	_ 110
		345		_			350					355				
atg	act	ggc	ctg	cct	ctg	tct	gct	ctt	cca	cca	cct	ctg	cac	aaa	gcc	1458
Met		Gly	Leu	Pro	Leu		Ala	Leu	Pro	Pro	Pro	Leu	His	Lys	Ala	
	360					365					370					

cag	tcc	tcc	ggc	cca	gaa	cat	cct	ggc	ccg	gag	tcc	tcc	ctg	ccc	tca	1506	
Gln	Ser	Ser	Gly	Pro	Glu	His	Pro	Gly	Pro	Glu	Ser	Ser	Leu	Pro	Ser		
375					380					385					390		
ggg	gct	ctc	agc	aag	tca	gct	cct	ggg	tcc	ttc	tgg	cac	att	cag	gca	1554	
Gly	Ala	Leu	Ser	Lys	Ser	Ala	Pro	Gly	Ser	Phe	Trp	His	Ile	Gln	Ala		
				395					400					405			
gat	cat	gca	tac	cag	gct	ctg	cca	tcc	ttc	cag	atc	cca	gtc	tca	cca	1602	
Asp	His	Ala	Tyr	Gln	Ala	Leu	Pro	Ser	Phe	Gln	Ile	Pro	Val	Ser	Pro		
			410					415					420				
cac	atc	tac	acc	agt	gtc	agc	tgg	gct	gct	gcc	ccc	tcc	gcc	gcc	tgc	1650	
His	Ile	Tyr	Thr	Ser	Val	Ser	Trp	Ala	Ala	Ala	Pro	Ser	Ala	Ala	Cys		
		425					430					435					
tct	ctc	tct	ccg	gtc	cgg	agc	cgg	tcg	cta	agc	ttc	agc	gag	CCC	cag	1698	
Ser	Leu	Ser	Pro	Val	Arg	Ser	Arg	Ser	Leu	Ser	Phe	Ser	Glu	Pro	Gln		
	440					445					450						
cag	cca	gca	cct	gcg	atg	aaa	tct	cat	ctg	atc	gtc	act	tct	cca	ccc	1746	
Gln	Pro	Ala	Pro	Ala	Met	Lys	Ser	His	Leu	Ile	Val	Thr	Ser	Pro	Pro		
455					460					465					470		
cgg	gcc	cag	agt	ggt	gcc	agg	aaa	gcc	cga	ggg	gag	gct	aag	aag	tgc	1794	
Arg	Ala	Gln	Ser	Gly	Ala	Arg	Lys	Ala	Arg	Gly	Glu	Ala	Lys	Lys	Cys		
				475					480					485			
cgc	aag	gtg	tat	ggc	atc	gag	cac	cgg	gac	cag	tgg	tgc	acg	gcg	tgc	1842	
Arg	Lys	Val	Tyr	Gly	Ile	Glu	His	Arg	Asp	Gln	Trp	Cys	Thr	Ala	Cys		
			490					495					500				
cgg	tgg	aag	aag	gcc	tgc	cag	cgc	ttt	ctg	gac	tga	gct	gtgci	tgc		1888	
Arg	Trp	Lys	Lys	Ala	Суз	Gln	Arg	Phe	Leu	Asp							
		505					510										
aggttctact ctgttcctgg ccctgccggc agccactgac										gac	e aagaggecag tgtgteacea 1						
gccctcagca gaaaccgaaa gagaaagaac								c gga	aaaca	acgg	agti	tgg	gct (	ctgtt	ggcta	2008	

<210> 2 <211> 513 <212> PRT <213> Homo sapiens

<400> 2

Met	Ala	Ser	Val	Leu	Ser	Arg	Arg	Leu	Gly	Lys	Arg	Ser	Leu	Leu	Gly
1				5					10					15	
Ala	Arg	Val	Leu	Gly	Pro	Ser	Ala	Ser	Glu	Gly	Pro	Ser	Ala	Ala	Pro
			20					25					30		
Pro	Ser	Glu	Pro	Leu	Leu	Glu	Gly	Ala	Ala	Pro	Gln	Pro	Phe	Thr	Thr
		35					40					45			
Ser	Asp	Asp	Thr	Pro	Суз	Gln	Glu	Gln	Pro	Lys	Glu	Val	Leu	Lys	Ala
	50					55					60				
Pro	Ser	Thr	Ser	Gly	Leu	Gln	Gln	Val	Ala	Phe	Gln	Pro	Gly	Gln	Lys
65					70					75					80
Val	Tyr	Val	Trp	Tyr	Gly	Gly	Gln	Glu	Суѕ	Thr	Gly	Leu	Val	Glu	Gln
				85					90					95	
His	Ser	Trp	Met	Glu	Gly	Gln	Val	Thr	Val	Trp	Leu	Leu	Glu	Gln	Lys
			100					105					110		
Leu	Gln	Val	Суз	Суз	Arg	Val	Glu	Glu	Val	Trp	Leu	Ala	Glu	Leu	Gln
		115					120					125			
Gly	Pro	Cys	Pro	Gln	Ala	Pro	Pro	Leu	Glu	Pro	Gly	Ala	Gln	Ala	Leu
	130					135					140				
Ala	Tyr	Arg	Pro	Val	Ser	Arg	Asn	Ile	Asp	Val	Pro	Lys	Arg	Lys	Ser

155 145 150 Asp Ala Val Glu Met Asp Glu Met Met Ala Ala Met Val Leu Thr Ser 165 170 Leu Ser Cys Ser Pro Val Val Gln Ser Pro Pro Gly Thr Glu Ala Asn 185 Phe Ser Ala Ser Arg Ala Ala Cys Asp Pro Trp Lys Glu Ser Gly Asp 200 Ile Ser Asp Ser Gly Ser Ser Thr Thr Ser Gly His Trp Ser Gly Ser 215 220 Ser Gly Val Ser Thr Pro Ser Pro Pro His Pro Gln Ala Ser Pro Lys 235 230 Tyr Leu Gly Asp Ala Phe Gly Ser Pro Gln Thr Asp His Gly Phe Glu 245 250 Thr Asp Pro Asp Pro Phe Leu Leu Asp Glu Pro Ala Pro Arg Lys Arg 265 Lys Asn Ser Val Lys Val Met Tyr Lys Cys Leu Trp Pro Asn Cys Gly 280 Lys Val Leu Arg Ser Ile Val Gly Ile Lys Arg His Val Lys Ala Leu 295 His Leu Gly Asp Thr Val Asp Ser Asp Gln Phe Lys Arg Glu Glu Asp 310 Phe Tyr Tyr Thr Glu Val Gln Leu Lys Glu Glu Ser Ala Ala Ala Ala 330 325 Ala Ala Ala Ala Gly Thr Pro Val Pro Gly Thr Pro Thr Ser Glu 345 Pro Ala Pro Thr Pro Ser Met Thr Gly Leu Pro Leu Ser Ala Leu Pro 360 Pro Pro Leu His Lys Ala Gln Ser Ser Gly Pro Glu His Pro Gly Pro 375 Glu Ser Ser Leu Pro Ser Gly Ala Leu Ser Lys Ser Ala Pro Gly Ser 390 395 Phe Trp His Ile Gln Ala Asp His Ala Tyr Gln Ala Leu Pro Ser Phe 405 410 Gln Ile Pro Val Ser Pro His Ile Tyr Thr Ser Val Ser Trp Ala Ala 425 420 Ala Pro Ser Ala Ala Cys Ser Leu Ser Pro Val Arg Ser Arg Ser Leu 440 Ser Phe Ser Glu Pro Gln Gln Pro Ala Pro Ala Met Lys Ser His Leu 455 460 Ile Val Thr Ser Pro Pro Arg Ala Gln Ser Gly Ala Arg Lys Ala Arg 470 475 Gly Glu Ala Lys Lys Cys Arg Lys Val Tyr Gly Ile Glu His Arg Asp 485 490 Gln Trp Cys Thr Ala Cys Arg Trp Lys Lys Ala Cys Gln Arg Phe Leu 500 505 Asp

<210> 3 <211> 1901 <212> DNA <213> Homo sapiens <220>

<221> misc_feature <222> (1191)..(1191) <400> 3 ggctggcaga gctgcagggc ccctgtcccc aggcaccacc cctggagccc ggagcccagg 60 ccctggccta caggcccgtc tccaggaaca tcgatgtccc aaagaggaag tcggacgcag 120 tggaaatgga tgagatgatg gcggccatgg tgctgacgtc cctgtcctgc agccctgttg 180 tacagagtcc tcccgggacc gaggccaact tctctgcttc ccgtgcggcc tgcgacccat 240 ggaaggagag tggtgacatc tcggacagcg gcagcagcac taccagcggt cactggagtg 300 ggagcagtgg tgtctccacc ccctcgcccc cccaccccca ggccagcccc aagtatttgg 360 gggatgettt tggtteteec caaactgate atggetttga gaccgateet gaccetttee 420 tgctggacga accagctcca cgaaaaagaa agaactctgt gaaggtgatg tacaagtgcc 480 tgtggccaaa ctgtggcaaa gttctgcgct ccattgtggg catcaaacga cacgtcaaag 540 ccctccatct gggggacaca gtggactctg atcagttcaa gcgggaggag gatttctact 600 acacagaggt gcagctgaag gaggaatctg ctgctgctgc tgctgctgct gccgcaggca 660 ccccagtccc tgggactccc acctccgage cagetcccac ccccagcatg actggcctgc 720 ctctgtctgc tcttccacca cctctgcaca aagcccagtc ctccggccca gaacatcctg 780 geoeggagte etecetgeee teaggggete teageaagte ageteetggg teettetgge 840 acatteagge agateatgea taccaggete tgecateett ceagateeca gteteaceae 900 acatctacac cagtgtcagc tgggctgctg ccccctccgc cgcctgctct ctctctccgg 960 teeggageeg gtegetaage tteagegage eecageagee ageacetgeg atgaaatete 1020 atctgatcgt cactteteca eeeegggeee agagtggtge eaggtgagat gteegetgte 1080 gtcccctgcc ttctggtttc tgtgccctgt ctccagtggc gtggactccg accccaccca 1140 gatgaagtca ccagggttag tccccagaga ggagcccaga tggcggatgc nccagatggg 1200 atgactgttt ggtcctcaga gcctctggcc cctggtcctg gtgacttttg ccgggagctg 1260 cccctttggc ctctgcttgt tctcccagcc ccacttggcc actctcctgg gcccaccacc 1320 tgtgtggggc tcgatttgca ttcctctctt tctgcaggaa agcccgaggg gaggctaaga 1380 agtgccgcaa ggtgtatggc atcgagcacc gggaccagtg gtgcacggcg tgccggtgga 1440 agaaggcctg ccagcgcttt ctggactgag ctgtgctgca ggttctactc tgttcctggc 1500 cctgccggca gccactgaca agaggccagt gtgtcaccag ccctcagcag aaaccgaaag 1560 agaaagaacg gaaacacgga gtttgggctc tgttggctaa ggtgtaacac ttaaagcaat 1620 tttctcccat tgtgcgaaca ttttattttt taaaaaaaag aaacaaaaat atttttcccc 1680 ctaaaatagg agagagccaa aactgaccaa ggctattcag cagtgaacca gtgaccaaag 1740 aattaattac cctccgtttc ccacatcccc actctctagg ggattagctt gtgcgtgtca 1800 aaagaaggaa cagctcgttc tgcttcctgc tgagtcggtg aattctttgc tttctaaact 1860 1901 cttccagaaa ggactgtgag caagatgaat ttacttttct t <210> 4 <211> 33 <212> DNA <213> Artificial Sequence <220> <223> PCR primer used to amplify PBF cDNA <400> 4 33 tactagctag ctaaggcagc agcatggcga gtg <210> 5 <211> 34 <212> DNA <213> Artificial Sequence <220>

<223> PCR primer used to amplify PBF cDNA

aaatatgcgg ccgcggccag gaacagagta gaac

<400> 5

```
<210> 6
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic peptide derived from SEQ ID NO: 2 that can
      be recognized by CTLs when bound to HLA-A24 antigen
Ala Tyr Arg Pro Val Ser Arg Asn Ile
                  5
<210> 7
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic peptide derived from SEQ ID NO: 2 that can
      be recognized by CTLs when bound to HLA-A24 antigen
<400> 7
Asp Phe Tyr Tyr Thr Glu Val Gln Leu
<210> 8
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic peptide derived from SEQ ID NO: 2 that can
      be recognized by CTLs when bound to \ensuremath{\mathsf{HLA}}\xspace-\ensuremath{\mathsf{A24}}\xspace antigen
<400> 8
Gly Phe Glu Thr Asp Pro Asp Pro Phe
 1
                   5
<210> 9
<211> 9
<212> PRT
<213> Artificial Sequence
<223> Synthetic peptide derived from SEQ ID NO: 2 that can
      be recognized by CTLs when bound to HLA-A24 antigen
<400> 9
```

Lys Tyr Leu Gly Asp Ala Phe Gly Ser

```
5
 1
<210> 10
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic peptide derived from SEQ ID NO: 2 that can
      be recognized by CTLs when bound to HLA-A24 antigen
<400> 10
Arg Ser Leu Leu Gly Ala Arg Val Leu
<210> 11
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic peptide derived from SEQ ID NO: 2 that can
      be recognized by CTLs when bound to HLA-A24 antigen
<400> 11
Ala Ala Pro Pro Ser Glu Pro Leu Leu
 1
                  5
<210> 12
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic peptide derived from SEQ ID NO: 2 that can
      be recognized by CTLs when bound to HLA-A24 antigen
<400> 12
Ile Tyr Thr Ser Val Ser Trp Ala Ala
<210> 13
<211> 9
<212> PRT
<213> Artificial Sequence
```

<220>

<223> Synthetic peptide derived from SEQ ID NO: 2 that can be recognized by CTLs when bound to HLA-A24 antigen

```
<210> 14
<211> 9
<212> PRT
<213> Artificial Sequence
<220>
<223> Synthetic peptide derived from SEQ ID NO: 2 that can be recognized by CTLs when bound to HLA-A24 antigen
```

<400> 14